

Art Unit: 1652

APPENDIX A

RESULT 2

US-09-946-374-85

; Sequence 85, Application US/09946374

; Publication No. US20030073129A1

; GENERAL INFORMATION:

; APPLICANT: Baker, Kevin P.

; APPLICANT: Botstein, David

; APPLICANT: Desnoyers, Luc

; APPLICANT: Eaton, Dan L.

; APPLICANT: Ferrara, Napoleone

; APPLICANT: Fong, Sherman

; APPLICANT: Gao, Wei-Qiang

; APPLICANT: Goddard, Audrey

; APPLICANT: Godowski, Paul J.

; APPLICANT: Grimaldi, Christopher J.

; APPLICANT: Gurney, Austin L.

; APPLICANT: Hillan, Kenneth J.

; APPLICANT: Pan, James

; APPLICANT: Paoni, Nicholas F.

; APPLICANT: Roy, Margaret Ann

; APPLICANT: Smith, Victoria

; APPLICANT: Stewart, Timothy A.

; APPLICANT: Tumas, Daniel

; APPLICANT: Watanabe, Colin K.

; APPLICANT: Williams, P. Mickey

; APPLICANT: Wood, William I.

; TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

; TITLE OF INVENTION: Acids Encoding the Same

; FILE REFERENCE: P2830P1C1

; CURRENT APPLICATION NUMBER: US/09/946,374

; CURRENT FILING DATE: 2001-09-04

; PRIOR APPLICATION NUMBER: 60/098716

; PRIOR FILING DATE: 1998-09-01

; PRIOR APPLICATION NUMBER: 60/098723

; PRIOR FILING DATE: 1998-09-01

; PRIOR APPLICATION NUMBER: 60/098749

; PRIOR FILING DATE: 1998-09-01

; PRIOR APPLICATION NUMBER: 60/098750

; PRIOR FILING DATE: 1998-09-01

; PRIOR APPLICATION NUMBER: 60/098803

; PRIOR FILING DATE: 1998-09-02

; PRIOR APPLICATION NUMBER: 60/098821

; PRIOR FILING DATE: 1998-09-02

; PRIOR APPLICATION NUMBER: 60/098843

; PRIOR FILING DATE: 1998-09-02

; PRIOR APPLICATION NUMBER: 60/099536

; PRIOR FILING DATE: 1998-09-09

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; PRIOR FILING DATE: 1998-09-09

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; PRIOR FILING DATE: 1998-09-09

; PRIOR APPLICATION NUMBER: 60/099741

; PRIOR FILING DATE: 1998-09-10

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; PRIOR APPLICATION NUMBER: 60/101741
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; PRIOR FILING DATE: 1998-09-24

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; PRIOR APPLICATION NUMBER: 60/101915
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; PRIOR FILING DATE: 1998-10-26
; PRIOR APPLICATION NUMBER: 60/105694
; PRIOR FILING DATE: 1998-10-26
; PRIOR APPLICATION NUMBER: 60/105807

Query Match 98.3%; Score 3261; DB 10; Length 3316;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 3311; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY	1	CATGGCCTGACTCGGGACAGCTCAGAGCAGGGCAGAACTGGGGACACTCTGGGCCGGCCT	60
Db	5	CATGGCCTGACTCGGGACAGCTCAGAGCAGGGCAGAACTGGGGACACTCTGGGCCGGCCT	64
QY	61	TCTGCCTGCATGGACGCTCTGAAGCCACCCTGTCTCTGGAGGAACCACGAGCGAGGGAAG	120
Db	65	TCTGCCTGCATGGACGCTCTGAAGCCACCCTGTCTCTGGAGGAACCACGAGCGAGGGAAG	124
QY	121	AAGGACAGGGACTCGTGTGGCAGGAAGAACTCAGAGCCGGGAAGCCCCATTCACTAGAA	180
Db	125	AAGGACAGGGACTCGTGTGGCAGGAAGAACTCAGAGCCGGGAAGCCCCATTCACTAGAA	184
QY	181	GCACTGAGAGATGCGGCCCCCTCGCAGGGTCTGAATTTCTGCTGCTGTTACAAAGATG	240
Db	185	GCACTGAGAGATGCGGCCCCCTCGCAGGGTCTGAATTTCTGCTGCTGTTACAAAGATG	244
QY	241	CTTTTTATCTTTAACTTTTTGTTTTCCCCACTTCCGACCCCGGCGTTGATCTGCATCCTG	300
Db	245	CTTTTTATCTTTAACTTTTTGTTTTCCCCACTTCCGACCCCGGCGTTGATCTGCATCCTG	304
QY	301	ACATTTGGAGCTGCCATCTTCTTGTGGCTGATCACCAGACCTCAACCCGTCTTACCTCTT	360
Db	305	ACATTTGGAGCTGCCATCTTCTTGTGGCTGATCACCAGACCTCAACCCGTCTTACCTCTT	364
QY	361	CTTGACCTGAACAATCAGTCTGTGGGAATTGAGGGAGGAGCACGGAAGGGGGTTTCCCAG	420
Db	365	CTTGACCTGAACAATCAGTCTGTGGGAATTGAGGGAGGAGCACGGAAGGGGGTTTCCCAG	424
QY	421	AAGAACAATGACCTAACAAGTTGCTGCTTCTCAGATGCCAAGACTATGTATGAGGTTTTC	480
Db	425	AAGAACAATGACCTAACAAGTTGCTGCTTCTCAGATGCCAAGACTATGTATGAGGTTTTC	484
QY	481	CAAAGAGGACTCGCTGTGTCTGACAATGGGCCCTGCTTGGGATATAGAAAACCAAACCAG	540
Db	485	CAAAGAGGACTCGCTGTGTCTGACAATGGGCCCTGCTTGGGATATAGAAAACCAAACCAG	544
QY	541	CCCTACAGATGGCTATCTTACAAACAGGTGTCTGATAGAGCAGAGTACCTGGGTTCTGT	600
Db	545	CCCTACAGATGGCTATCTTACAAACAGGTGTCTGATAGAGCAGAGTACCTGGGTTCTGT	604
QY	601	CTCTTGATATAAGGTTATAAATCATCACCAGACCAGTTTGTTCGGCATCTTTGCTCAGAAT	660
Db	605	CTCTTGATATAAGGTTATAAATCATCACCAGACCAGTTTGTTCGGCATCTTTGCTCAGAAT	664
QY	661	AGGCCAGAGTGGATCATCTCCGAATTGGCTTGTTACACGTAATCTATGGTAGCTGTACCT	720
Db	665	AGGCCAGAGTGGATCATCTCCGAATTGGCTTGTTACACGTAATCTATGGTAGCTGTACCT	724
QY	721	CTGTATGACACCTTGGGACCAGAAGCCATCGTACATATTGTCAACAAGGCTGATATCGCC	780
Db	725	CTGTATGACACCTTGGGACCAGAAGCCATCGTACATATTGTCAACAAGGCTGATATCGCC	784
QY	781	GTGGTGATCTGTGACACACCCCAAAAGGCATTGGTGCTGATAGGGAATGTAGAGAAAGGC	840
Db	785	ATGGTGATCTGTGACACACCCCAAAAGGCATTGGTGCTGATAGGGAATGTAGAGAAAGGC	844
QY	841	TTACCCCCGAGCCTGAAGGTGATCATCCTTATGGACCCCTTTGATGATGACCTGAAGCAA	900

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Db 845 TTCACCCCGAGCCTGAAGGTGATCATCCTTATGGACCCCTTTGATGATGACCTGAAGCAA 904

Qy 901 AGAGGGGAGAAGAGTGGAAATTGAGATCTTATCCCTATATGATGCTGAGAACCTAGGCAAA 960
|||||

Db 905 AGAGGGGAGAAGAGTGGAAATTGAGATCTTATCCCTATATGATGCTGAGAACCTAGGCAAA 964

Qy 961 GAGCACTTCAGAAAACCTGTGCCTCCTAGCCCAGAAGACCTGAGCGTCATCTGCTTCACC 1020
|||||

Db 965 GAGCACTTCAGAAAACCTGTGCCTCCTAGCCCAGAAGACCTGAGCGTCATCTGCTTCACC 1024

Qy 1021 AGTGGGACCACAGGTGACCCCAAAGGAGCCATGATAACCCATCAAAATATTGTTTCAAAT 1080
|||||

Db 1025 AGTGGGACCACAGGTGACCCCAAAGGAGCCATGATAACCCATCAAAATATTGTTTCAAAT 1084

Qy 1081 GCTGCTGCCTTTTCTCAAATGTGTGGAGCATGCTTATGAGCCCACTCCTGATGATGTGGCC 1140
|||||

Db 1085 GCTGCTGCCTTTTCTCAAATGTGTGGAGCATGCTTATGAGCCCACTCCTGATGATGTGGCC 1144

Qy 1141 ATATCCTACCTCCCTCTGGCTCATATGTTTGAGAGGATTGTACAGGCTGTTGTGTACAGC 1200
|||||

Db 1145 ATATCCTACCTCCCTCTGGCTCATATGTTTGAGAGGATTGTACAGGCTGTTGTGTACAGC 1204

Qy 1201 TGTGGAGCCAGAGTTGGATTCTTCCAAGGGGATATTCGGTTGCTGGCTGACGACATGAAG 1260
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Db 1205 TGTGGAGCCAGAGTTGGATTCTTCCAAGGGGATATTCGGTTGCTGGCTGACGACATGAAG 1264

Qy 1261 ACTTTGAAGCCACATTGTTTCCCGCGGTGCCTCGACTCCTTAACAGGATCTACGATAAG 1320
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Db 1265 ACTTTGAAGCCACATTGTTTCCCGCGGTGCCTCGACTCCTTAACAGGATCTACGATAAG 1324

Qy 1321 GTACAAAATGAGGCCAAGACACCCTTGAAGAAGTTCTTGTGAAGCTGGCTGTTTCCAGT 1380
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Db 1325 GTACAAAATGAGGCCAAGACACCCTTGAAGAAGTTCTTGTGAAGCTGGCTGTTTCCAGT 1384

Qy 1381 AAATTCAAAGAGCTTCAAAGGGTATCATCAGGCATGATAGTTTCTGGGACAAGCTCATC 1440
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Db 1385 AAATTCAAAGAGCTTCAAAGGGTATCATCAGGCATGATAGTTTCTGGGACAAGCTCATC 1444

Qy 1441 TTTGCAAAGATCCAGGACAGCCTGGGCGGAAGGGTTTCGTGTAATTGTCACTGGAGCTGCC 1500
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Db 1445 TTTGCAAAGATCCAGGACAGCCTGGGCGGAAGGGTTTCGTGTAATTGTCACTGGAGCTGCC 1504

Qy 1501 CCCATGTCCACTTCAGTCATGACATTCTTCCGGGCAGCAATGGGATGTCAGGTGTATGAA 1560
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Db 1505 CCCATGTCCACTTCAGTCATGACATTCTTCCGGGCAGCAATGGGATGTCAGGTGTATGAA 1564

Qy 1561 GCTTATGGTCAAACAGAATGCACAGGTGGCTGTACATTTACATTACCTGGGGACTGGACA 1620
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Db 1565 GCTTATGGTCAAACAGAATGCACAGGTGGCTGTACATTTACATTACCTGGGGACTGGACA 1624

Qy 1621 TCAGGTCACGTTGGGGTGCCCCCTGGCTTGCAATTACGTGAAGCTGGAAGATGTGGCTGAC 1680
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Db 1625 TCAGGTCACGTTGGGGTGCCCCCTGGCTTGCAATTACGTGAAGCTGGAAGATGTGGCTGAC 1684

Qy 1681 ATGAACTACTTTACAGTGAATAATGAAGGAGAGGTCTGCATCAAGGGTACAAACGTGTTC 1740
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Db 1685 ATGAACTACTTTACAGTGAATAATGAAGGAGAGGTCTGCATCAAGGGTACAAACGTGTTC 1744

Qy 1741 AAAGGATACCTGAAGGACCCTGAGAAGACACAGGAAGCCCTGGACAGTGATGGCTGGCTT 1800
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Db 1745 AAAGGATACCTGAAGGACCCTGAGAAGACACAGGAAGCCCTGGACAGTGATGGCTGGCTT 1804

Qy 1801 CACACAGGAGACATTGGTTCGCTGGCTCCCGAATGGAACCTCTGAAGATCATCGACCGTAAA 1860
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Db 1805 CACACAGGAGACATTGGTTCGCTGGCTCCCGAATGGAACCTCTGAAGATCATCGACCGTAAA 1864

Qy 1861 AAGAACATTTTCAAGCTGGCCCAAGGAGAATACATTGCACCAGAGAAGATAGAAAATATC 1920

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Db 1865 AAGAACATTTTCAAGCTGGCCCAAGGAGAATACATTGCACCAGAGAAGATAGAAAATATC 1924
Qy 1921 TACAACAGGAGTCAACCAGTGTTACAAATTTTGTACACGGGGAGAGCTTACGGTCATCC 1980
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Db 1925 TACAACAGGAGTCAACCAGTGTTACAAATTTTGTACACGGGGAGAGCTTACGGTCATCC 1984
Qy 1981 TTAGTAGGAGTGGTGGTTCCTGACACAGATGTACTTCCCTCATTTGCAGCCAAGCTTGGG 2040
|||||
Db 1985 TTAGTAGGAGTGGTGGTTCCTGACACAGATGTACTTCCCTCATTTGCAGCCAAGCTTGGG 2044
Qy 2041 GTGAAGGGCTCCTTTGAGGAACTGTGCCAAAACCAAGTTGTAAGGGAAGCCATTTTAGAA 2100
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Db 2045 GTGAAGGGCTCCTTTGAGGAACTGTGCCAAAACCAAGTTGTAAGGGAAGCCATTTTAGAA 2104
Qy 2101 GACTTGCAGAAAATTGGGAAAGAAAGTGGCCTTAAACTTTTGAACAGGTCAAAGCCATT 2160
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Db 2105 GACTTGCAGAAAATTGGGAAAGAAAGTGGCCTTAAACTTTTGAACAGGTCAAAGCCATT 2164
Qy 2161 TTTCTTCATCCAGAGCCATTTTCCATTGAAAATGGGCTCTTGACACCAACATTGAAAGCA 2220
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Db 2165 TTTCTTCATCCAGAGCCATTTTCCATTGAAAATGGGCTCTTGACACCAACATTGAAAGCA 2224
Qy 2221 AAGCGAGGAGAGCTTTCCAAATACTTTTCGACCCAAATTGACAGCCTGTATGAGCACATC 2280
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Db 2225 AAGCGAGGAGAGCTTTCCAAATACTTTTCGACCCAAATTGACAGCCTGTATGAGCACATC 2284
Qy 2281 CAGGATTAGGATAAGGTACTTAAGTACCTGCCGGCCCACTGTGCACTGCTTGTGAGAAAA 2340
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Db 2285 CAGGATTAGGATAAGGTACTTAAGTACCTGCCGGCCCACTGTGCACTGCTTGTGAGAAAA 2344
Qy 2341 TGGATTAAAACTATTCTTACATTTGTTTTGCCTTTCCCTCTATTTTTTTTAACTGTT 2400
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Db 2345 TGGATTAAAACTATTCTTACATTTGTTTTGCCTTTCCCTCTATTTTTTTTAACTGTT 2404
Qy 2401 AAACCTAAAGCCATAGCTTTTGTGTTTATATTGAGACATATAATGTGTAACTTAGTTCC 2460
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Db 2405 AAACCTAAAGCCATAGCTTTTGTGTTTATATTGAGACATATAATGTGTAACTTAGTTCC 2464
Qy 2461 CAAATAAATCAATCCTGTCTTTCCCATCTTCGATGTTGCTAATATTAAGGCTTCAGGGCT 2520
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Db 2465 CAAATAAATCAATCCTGTCTTTCCCATCTTCGATGTTGCTAATATTAAGGCTTCAGGGCT 2524
Qy 2521 ACTTTTATCAACATGCCTGTCTTCAAGATCCCAGTTTATGTTCTGTGTCCTTCCTCATGA 2580
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Db 2525 ACTTTTATCAACATGCCTGTCTTCAAGATCCCAGTTTATGTTCTGTGTCCTTCCTCATGA 2584
Qy 2581 TTTCCAACCTTAATACTATTAGTAACCACAAGTTCAAGGGTCAAAGGGACCCTCTGTGCC 2640
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Db 2585 TTTCCAACCTTAATACTATTAGTAACCACAAGTTCAAGGGTCAAAGGGACCCTCTGTGCC 2644
Qy 2641 TTCTTCTTTGTTTTGTGATAAACATAAAGTCCCAACAGTCTCTATGCTTATTTACATCTT 2700
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Qy 2701 CTACTGTTCAAACCTAAGAGATTTTTAAATTCTGAAAACTGCTTACAATTCATGTTTTCT 2760
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Qy 2761 AGCCACTCCACAAACCACTAAAATTTTAGTTTTAGCCTATCACTCATGTCAATCATATCT 2820
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Db 2765 AGCCACTCCACAAACCACTAAAATTTTAGTTTTAGCCTATCACTCATGTCAATCATATCT 2824
Qy 2821 ATGAGACAAATGTCTCCGATGCTCTTCTGCGTAAATTAAATTGTGTACTGAAGGGAAAAG 2880
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Db 2825 ATGAGACAAATGTCTCCGATGCTCTTCTGCGTAAATTAAATTGTGTACTGAAGGGAAAAG 2884

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Qy	2881	TTTGATCATACCAAACATTTCCCTAAACTCTCTAGTTAGATATCTGACTTGGGAGTATTAA	2940
Db	2885	TTTGATCATACCAAACATTTCCCTAAACTCTCTAGTTAGATATCTGACTTGGGAGTATTAA	2944
Qy	2941	AAATTGGGTCTATGACATACTGTCCAAAAGGAATGCTGTTCTTAAAGCATTATTTACAGT	3000
Db	2945	AAATTGGGTCTATGACATACTGTCCAAAAGGAATGCTGTTCTTAAAGCATTATTTACAGT	3004
Qy	3001	AGGAACTGGGGAGTAAATCTGTTCCCTACAGTTTGCTGCTGAGCTGGAAGCTGTGGGGGA	3060
Db	3005	AGGAACTGGGGAGTAAATCTGTTCCCTACAGTTTGCTGCTGAGCTGGAAGCTGTGGGGGA	3064
Qy	3061	AGGAGTTGACAGGTGGGCCCAGTGAACCTTTTCCAGTAAATGAAGCAAGCACTGAATAAAA	3120
Db	3065	AGGAGTTGACAGGTGGGCCCAGTGAACCTTTTCCAGTAAATGAAGCAAGCACTGAATAAAA	3124
Qy	3121	ACCTCCTGAACTGGGAACAAAGATCTACAGGCAAGCAAGATGCCCACACAACAGGCTTAT	3180
Db	3125	ACCTCCTGAACTGGGAACAAAGATCTACAGGCAAGCAAGATGCCCACACAACAGGCTTAT	3184
Qy	3181	TTTCTGTGAAGGAACCAACTGATCTCCCCACCCTTGGATTAGAGTTCCTGCTCTACCTT	3240
Db	3185	TTTCTGTGAAGGAACCAACTGATCTCCCCACCCTTGGATTAGAGTTCCTGCTCTACCTT	3244
Qy	3241	ACCCACAGATAACACATGTTGTTTCTACTTGTAATGTAAAGTCTTTAAAATAAACTATT	3300
Db	3245	ACCCACAGATAACACATGTTGTTTCTACTTGTAATGTAAAGTCTTTAAAATAAACTATT	3304
Qy	3301	ACAGATAAAAAA	3312
Db	3305	ACAGATAAAAAA	3316